



SEQUENCE LISTING

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Neidhardt, Helge
Paulista, Michael

<120> NEW GROWTH/DIFFERENTIATING FACTOR OF TGF- FAMILY

<130> 100564-09022

<140> US 09/386,450

<141> 1999-08-31

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<151> 1994-08-10

<150> DE P 43 26 829.3

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<160> 41

<170> PatentIn version 3.0

<210> 1

<211> 2703

<212> DNA

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<222> (1)..(2703)

<223> coding region for TGF-beta protien MP-52

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<210> 2

<211> 501

<212> PRT

<213> Artificial/Unknown

<220>

<221> DOMAIN

<222> (1)..(501)

<223> TGF-beta protien MP-52

<400> 2

Met Arg Leu Pro Lys Leu Leu Thr Phe Leu Leu Trp Tyr Leu Ala Trp
1 5 10 15
Leu Asp Leu Glu Phe Ile Cys Thr Val Leu Gly Ala Pro Asp Leu Gly
20 25 30
Gln Arg Pro Gln Gly Thr Arg Pro Gly Leu Ala Lys Ala Glu Ala Lys
35 40 45
Glu Arg Pro Pro Leu Ala Arg Asn Val Phe Arg Pro Gly Gly His Ser
50 55 60
Tyr Gly Gly Gly Ala Thr Asn Ala Asn Ala Arg Ala Lys Gly Gly Thr
65 70 75 80
Gly Gln Thr Gly Gly Leu Thr Gln Pro Lys Lys Asp Glu Pro Lys Lys
85 90 95
Leu Pro Pro Arg Pro Gly Gly Pro Glu Pro Lys Pro Gly His Pro Pro
100 105 110
Gln Thr Arg Gln Ala Thr Ala Arg Thr Val Thr Pro Lys Gly Gln Leu
115 120 125
Pro Gly Gly Lys Ala Pro Pro Lys Ala Gly Ser Val Pro Ser Ser Phe
130 135 140
Leu Leu Lys Lys Ala Arg Glu Pro Gly Pro Pro Arg Glu Pro Lys Glu
145 150 155 160
Pro Phe Arg Pro Pro Pro Ile Thr Pro His Glu Tyr Met Leu Ser Leu
165 170 175
Tyr Arg Thr Leu Ser Asp Ala Asp Arg Lys Gly Gly Asn Ser Ser Val
180 185 190
Lys Leu Glu Ala Gly Leu Ala Asn Thr Ile Thr Ser Phe Ile Asp Lys
195 200 205
Gly Gln Asp Asp Arg Gly Pro Val Val Arg Lys Gln Arg Tyr Val Phe
210 215 220

Asp Ile Ser Ala Leu Glu Lys Asp Gly Leu Leu Gly Ala Glu Leu Arg
 225 230 235 240
 Ile Leu Arg Lys Lys Pro Ser Asp Thr Ala Lys Pro Ala Ala Pro Gly
 245 250 255
 Gly Gly Arg Ala Ala Gln Leu Lys Leu Ser Ser Cys Pro Ser Gly Arg
 260 265 270
 Gln Pro Ala Ser Leu Leu Asp Val Arg Ser Val Pro Gly Leu Asp Gly
 275 280 285
 Ser Gly Trp Glu Val Phe Asp Ile Trp Lys Leu Phe Arg Asn Phe Lys
 290 295 300
 Asn Ser Ala Gln Leu Cys Leu Glu Leu Glu Ala Trp Glu Arg Gly Arg
 305 310 315 320
 Ala Val Asp Leu Arg Gly Leu Gly Phe Asp Arg Ala Ala Arg Gln Val
 325 330 335
 His Glu Lys Ala Leu Phe Leu Val Phe Gly Arg Thr Lys Lys Arg Asp
 340 345 350
 Leu Phe Phe Asn Glu Ile Lys Ala Arg Ser Gly Gln Asp Asp Lys Thr
 355 360 365
 Val Tyr Glu Tyr Leu Phe Ser Gln Arg Arg Lys Arg Arg Ala Pro Leu
 370 375 380
 Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala Arg Cys
 385 390 395 400
 Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp Asp Asp
 405 410 415
 Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu Gly Leu
 420 425 430
 Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His Ala Val
 435 440 445
 Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro Pro Thr
 450 455 460
 Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe Ile Asp
 465 470 475 480
 Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val Val Glu
 485 490 495
 Ser Cys Gly Cys Arg
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<210> 3

<211> 24

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(24)

<223> MP-52 adapter primer

<400> 3
agaattcgca tgccatgggc gacg

24

<210> 4

<211> 23

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(23)

<223> MP-52 internal primer

<400> 4
cttgagtacg aggctttcca ctg

23

<210> 5

<211> 24

<212> DNA

<213> Artificial/Unknown

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<221> misc_feature

<222> (1)..(24)

<223> MP-52 adapter primer

<400> 5
attcgcatgc catggctcgac gaag

24

<210> 6

<211> 23

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(23)

<223> MP-52 internal primer

<400> 6
ggagcccacg aatcatgcag tca

23

<210> 7

<211> 23

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(23)

<223> MP-52 internal primer

<400> 7
acagcaggtg ggtggtgtgg act

23

<210> 8

<211> 44

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(44)

<223> adapter sequece

<400> 8
agaattcgca tgccatggtc gacgaagctt tttttttttt tttt

44

<210> 9

<211> 20

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(20)

<223> MP-52 internal primer

<400> 9
ccagcagccc atccttctcc

20

<210> 10

<211> 24

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(24)

<223> MP-52 internal primer

<400> 10
tccagggcac taatgtcaaa cacg

24

<210> 11

<211> 24

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(24)

<223> MP-52 internal primer

<400> 11
actaatgtca aacacgtacc tctg

24

<210> 12

<211> 10

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(10)

<223> adapter

<400> 12
agcggccgct

10

<210> 13

<211> 102

<212> PRT

<213> Artificial/Unknown

<220>

<221> DOMAIN

<222> (1)..(102)

<223> MP-52

<400> 13

Cys Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp Asp
1 5 10 15
Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu Gly
20 25 30
Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His Ala
35 40 45
Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro Pro
50 55 60
Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe Ile
65 70 75 80
Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val Val
85 90 95
Glu Ser Cys Gly Cys Arg
100

<210> 14

10

<211> 101

<212> PRT

<213> Artificial/Unknown

<220>

<221> DOMAIN

<222> (1)..(101)

<223> portion of BMP 2 corresponding to MP 52

<400> 14

Cys Lys Arg His Pro Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn
1 5 10 15

Asp Trp Ile Val Ala Pro Pro Gly Tyr His Ala Phe Tyr Cys His Gly
20 25 30

Glu Cys Pro Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala
35 40 45

Ile Val Gln Thr Leu Val Asn Ser Val Asn Ser Lys Ile Pro Lys Ala
50 55 60

Cys Cys Val Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp
65 70 75 80

Glu Asn Glu Lys Val Val Leu Lys Asn Tyr Gln Asp Met Val Val Glu
85 90 95

Gly Cys Gly Cys Arg
100

<210> 15

<211> 101

<212> PRT

<213> Artificial/Unknown

<220>

<221> DOMAIN

<222> (1)..(101)

<223> portion of BMP 4 corresponding to MP 52

<400> 15

Cys Arg Arg His Ser Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn
1 5 10 15
Asp Trp Ile Val Ala Pro Pro Gly Tyr Gln Ala Phe Tyr Cys His Gly
20 25 30
Asp Cys Pro Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala
35 40 45
Ile Val Gln Thr Leu Val Asn Ser Val Asn Ser Ser Ile Pro Lys Ala
50 55 60
Cys Cys Val Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp
65 70 75 80
Glu Tyr Asp Lys Val Val Leu Lys Asn Tyr Gln Glu Met Val Val Glu
85 90 95
Gly Cys Gly Cys Arg
100

<210> 16

<211> 102

<212> PRT

<213> Artificial/Unknown

<220>

<221> DOMAIN

<222> (1)..(102)

<223> portion of BMP 5 corresponding to MP 52

<400> 16

Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg Asp Leu Gly Trp Gln
1 5 10 15
Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala Phe Tyr Cys Asp Gly
20 25 30
Glu Cys Ser Phe Pro Leu Asn Ala His Met Asn Ala Thr Asn His Ala
12

35	40	45
Ile Val Gln Thr Leu Val His Leu Met Phe Pro Asp His Val Pro Lys		
50	55	60
Pro Cys Cys Ala Pro Thr Lys Leu Asn Ala Ile Ser Val Leu Tyr Phe		
65	70	75
Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr Arg Asn Met Val Val		
85	90	95
Arg Ser Cys Gly Cys His		
100		

<210> 17

<211> 102

<212> PRT

<213> Artificial/Unknown

<220>

<221> DOMAIN

<222> (1)..(102)

<223> portion of BMP 6 corresponding to MP 52

<400> 17

Cys Arg Lys His Glu Leu Tyr Val Ser Phe Gln Asp Leu Gly Trp Gln		
1	5	10
Asp Trp Ile Ile Ala Pro Lys Gly Tyr Ala Ala Asn Tyr Cys Asp Gly		
20	25	30
Glu Cys Ser Phe Pro Leu Asn Ala His Met Asn Ala Thr Asn His Ala		
35	40	45
Ile Val Gln Thr Leu Val His Leu Met Asn Pro Glu Tyr Val Pro Lys		
50	55	60
Pro Cys Cys Ala Pro Thr Lys Leu Asn Ala Ile Ser Val Leu Tyr Phe		
65	70	75
Asp Asp Asn Ser Asn Val Ile Leu Lys Lys Tyr Arg Asn Met Val Val		
85	90	95
Arg Ala Cys Gly Cys His		
100		

<210> 18

<211> 102

<212> PRT

<213> Artificial/Unknown

<220>

<221> DOMAIN

<222> (1)..(102)

<223> portion of BMP 7 corresponding to MP 52

<400> 18

Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg Asp Leu Gly Trp Gln
1 5 10 15

Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala Tyr Tyr Cys Glu Gly
20 25 30

Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn Ala Thr Asn His Ala
35 40 45

Ile Val Gln Thr Leu Val His Phe Ile Asn Pro Glu Thr Val Pro Lys
50 55 60

Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile Ser Val Leu Tyr Phe
65 70 75 80

Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr Arg Asn Met Val Val
85 90 95

Arg Ala Cys Gly Cys His
100

<210> 19

<211> 36

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(36)

<223> primer OD

<400> 19
atgaattccc atggacctgg gctggmakga mtggat

36

<210> 20

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of BMP 2 corresponding to primer OD

<400> 20
acgtgggggtg gaatgactgg at

22

<210> 21

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of BMP 3 corresponding to primer OD

<400> 21
atattggctg gagtgaatgg at

22

<210> 22

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of BMP 4 corresponding to primer OD

<400> 22

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22

<210> 23

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of BMP 7 corresponding to primer OD

<400> 23

acctgggctg gcaggactgg at

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<210> 24

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of TGF-beta-1 corresponding to primer OD

<400> 24
aggacctcgg ctggaagtgg at

22

<210> 25

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of TGF-beta-2 corresponding to primer OD

<400> 25
gggatctagg gtggaaatgg at

22

<210> 26

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of TGF-beta-3 corresponding to primer OD

<400> 26
aggatctggg ctggaagtgg gt

22

<210> 27

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of Inhibin alpha corresponding to primer OD

<400> 27
agctgggctg ggaacggtgg at

22

<210> 28

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of Inhibin beta-gamma corresponding to primer OD

<400> 28
acatcggctg gaatgactgg at

22

<210> 29

<211> 22

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(22)

<223> portion of Inhibin beta-beta corresponding to primer OD

<400> 29
tcatcggctg gaacgactgg at

22

<210> 30

<211> 29

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(29)

<223> Primer OID

<400> 30
atgaattcga gctgcgtsgg srcacagca

29

<210> 31

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of BMP 2 corresponding to primer OID

<400> 31
gagttctgtc gggacacagc a

21

<210> 32

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of BMP 3 corresponding to primer OID

<400> 32
catcttttct ggtacacagc a

21

<210> 33

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of BMP 4 corresponding to primer OID

<400> 33
cagttcagtg ggcacacaac a

21

<210> 34

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of BMP 7 corresponding to primer OID

<400> 34
gagctgcgtg ggcgcacagc a

21

<210> 35

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of TGF-beta-1 corresponding to primer OID

<400> 35
cagcgcctgc ggcacgcagc a

21

<210> 36

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of TGF-beta-2 corresponding to primer OID

<400> 36
taaatcttgg gacacgcagc a

21

<210> 37

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of TGF-beta-3 corresponding to primer OID

<400> 37
caggtcctgg ggcacgcagc a

21

<210> 38

<211> 21

<212> DNA

<213> Artificial/Unknown

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<221> misc_feature

<222> (1)..(21)

<223> portion of Inhibin alpha corresponding to primer OID

<400> 38

ccctgggaga gcagcacagc a

21

<210> 39

<211> 21

<212> DNA

<213> Artificial/Unknown

<220>

<221> misc_feature

<222> (1)..(21)

<223> portion of Inhibin beta-gamma corresponding to primer OID

<400> 39

cagcttggtg ggcacacagc a

21

<210> 40

<211> 21

<212> DNA

<213> Artificial/Unknown

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<222> (1)..(21)

<223> portion of Inhibin beta-beta corresponding to primer OID

<400> 40
cagcttggtg ggaatgcagc a

21

<210> 41

<211> 10

<212> DNA

<213> Artificial/Unknown

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<221> misc_feature

<222> (1)..(10)

<223> Adapter

<400> 41
tcgccggcga

10